



ENTERED

1636

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/441,242A

DATE: 02/26/2002

TIME: 09:30:11

#13

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Output Set: N:\CRF3\02262002\I441242A.raw

3 <110> APPLICANT: Russo, Giandomenico
 4 Croce, Carlo
 6 <120> TITLE OF INVENTION: TCL-1 Gene and Protein and Related Methods and Compositions
 8 <130> FILE REFERENCE: 8666-008
 10 <140> CURRENT APPLICATION NUMBER: 09/441,242A
 11 <141> CURRENT FILING DATE: 1999-11-16
 13 <150> PRIOR APPLICATION NUMBER: 08/330,272
 14 <151> PRIOR FILING DATE: 1994-10-07
 16 <160> NUMBER OF SEQ ID NOS: 12
 18 <170> SOFTWARE: PatentIn version 3.0
 20 <210> SEQ ID NO: 1
 21 <211> LENGTH: 1324
 22 <212> TYPE: DNA
 C--> 23 <213> ORGANISM: Artificial
 25 <220> FEATURE:
 26 <223> OTHER INFORMATION: Description of Artificial Sequence: cDNA sequence of TCL-1
 28 <221> NAME/KEY: CDS
 29 <222> LOCATION: (49)..(387)
 31 <400> SEQUENCE: 1
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 34 1
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 36 Pro Thr Leu Gly Glu Ala Val Thr Asp His Pro Asp Arg Leu Trp Ala
 37 5 10 15
 38 tgg gag aag ttc gtg tat ttg gac gag aag cag cac gcc tgg ctg ccc 153
 39 Trp Glu Lys Phe Val Tyr Leu Asp Glu Lys Gln His Ala Trp Leu Pro
 40 20 25 30 35
 41 tta acc atc gag ata aag gat agg tta cag tta cgg gtg ctc ttg cgt 201
 42 Leu Thr Ile Glu Ile Lys Asp Arg Leu Gln Leu Arg Val Leu Leu Arg
 43 40 45 50
 44 cgg gaa gac gtc gtc ctg ggg agg cct atg acc ccc acc cag ata ggc 249
 45 Arg Glu Asp Val Val Leu Gly Arg Pro Met Thr Pro Thr Gln Ile Gly
 46 55 60 65
 47 cca agc ctg ctg cct atc atg tgg cag ctc tac cct gat gga cga tac 297
 48 Pro Ser Leu Leu Pro Ile Met Trp Gln Leu Tyr Pro Asp Gly Arg Tyr
 49 70 75 80
 50 cga tcc tca gac tcc agt ttc tgg cgc tta gtg tac cac atc aag att 345
 51 Arg Ser Ser Asp Ser Ser Phe Trp Arg Leu Val Tyr His Ile Lys Ile
 52 85 90 95
 53 gac ggc gtg gag gac atg ctt ctc gag ctg ctg cca gat gac 387
 54 Asp Gly Val Glu Asp Met Leu Leu Glu Leu Leu Pro Asp Asp
 55 100 105 110

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56 tgatgtatgg tcttggcagc acctgtctcc ttccacccca gggcctgagc ctggccagcc 447
57 tacaatgggg atgttgtgtt tctgttcacc ttcgtttact atgcctgtgt cttctccacc 507
58 acgctggggg ctgggaggaa tggacagaca gaggatgagc tctacccagg gcctgcagga 567
59 cctgcctgta gccactctg ctcgccttag cactaccact cctgccaagg aggattccat 627
60 ttggcagagc ttcttccagg tgcccagcta tacctgtgcc tcggcttttc tcagctggat 687
61 gatggtcttc agcctctttc tgtcccttct gtccctcaca gcactagtat ttcattgttc 747
62 acacccactc agctccgtga acttgtgaga acacagccga ttcacctgag caggacctct 807
63 gaaaccctgg accagtggtc tcacatgggtg ctacgcctgc atgtaaacac gcctgcaaac 867
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65 tgctgcccc caccaggttca cgtgcagctc aaggaaaaggc ctgaaaggag cccttatctg 987
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68 atagaggttg ggtttccccc ctttatagat ggtcacgcac ctgggtgtta caaagtgtga 1167
69 tgtggcatga atactttttg taatgattga ttaaagtcaa gatagtttat ctaacttcgt 1227
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71 cccaaataaa aaatattcat ggaaaaaaaa aaaaaaa 1324

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75 <212> TYPE: PRT

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78 <220> FEATURE:

79 <223> OTHER INFORMATION: Description of Artificial Sequence: cDNA sequence of TCL-1

81 <400> SEQUENCE: 2

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83 1 5 10 15
85 Leu Trp Ala Trp Glu Lys Phe Val Tyr Leu Asp Glu Lys Gln His Ala
86 20 25 30
88 Trp Leu Pro Leu Thr Ile Glu Ile Lys Asp Arg Leu Gln Leu Arg Val
89 35 40 45
91 Leu Leu Arg Arg Glu Asp Val Val Leu Gly Arg Pro Met Thr Pro Thr
92 50 55 60
94 Gln Ile Gly Pro Ser Leu Leu Pro Ile Met Trp Gln Leu Tyr Pro Asp
95 65 70 75 80
97 Gly Arg Tyr Arg Ser Ser Asp Ser Ser Phe Trp Arg Leu Val Tyr His
98 85 90 95
100 Ile Lys Ile Asp Gly Val Glu Asp Met Leu Leu Glu Leu Leu Pro Asp
101 100 105 110
103 Asp

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106 <210> SEQ ID NO: 3

107 <211> LENGTH: 560

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112 <223> OTHER INFORMATION: Description of Artificial Sequence: genomic sequence of TCL-

114 <400> SEQUENCE: 3

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117 gggcggcggg cgggtgcgct gctggccggg gcctcgagga aggcgcgggc cagctggggc 180
118 cgggtctgcy ttcccaggag ctgccaccgt tccaggagagc aagtcaggcc gggacgttag 240

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120 gcctgcgcgg gaccctcaact tgccaccaag acccccacaaa accccgcccc atcctgcctt 360
121 acgccccgcc ccaaggtcgt tctcccgacc cgggggtccc ccccaagacc gtcctcccg 420
122 cccgccgctt ggtggcggcc gcatgctgcc cgatatataa gggtcggccc cacatccag 480
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132 <223> OTHER INFORMATION: Description of Artificial Sequence: MTCPl protein

134 <221> NAME/KEY: SITE

135 <222> LOCATION: 108

136 <223> OTHER INFORMATION: Xaa = any amino acid

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141 Gln Glu Gly Ile Tyr Arg Asp Glu Tyr Gln Arg Thr Trp Val Ala Val

142 20 25 30

143 Val Glu Glu Thr Ser Phe Leu Arg Ala Arg Val Gln Gln Ile Gln

144 35 40 45

145 Val Pro Leu Gly Asp Ala Ala Arg Pro Ser His Leu Leu Thr Ser Gln

146 50 55 60

147 Leu Pro Leu Met Trp Gln Leu Tyr Pro Glu Glu Arg Tyr Met Asp Asn

148 65 70 75 80

149 Asn Ser Arg Leu Trp Gln Ile Gln His His Leu Met Val Arg Gly Val

150 85 90 95

W--> 151 Gln Glu Leu Leu Leu Lys Leu Leu Pro Asp Asp Xaa

152 100 105

153 <210> SEQ ID NO: 5

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155 <212> TYPE: DNA

C--> 156 <213> ORGANISM: Artificial

158 <220> FEATURE:

159 <223> OTHER INFORMATION: Description of Artificial Sequence: genomic DNA of TCL-1

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163 2225, 2279, 2501, 2523, 2545, 2553, 2566, 2592, 2599, 3955, 3959,

164 3975, 3976, 3982, 3984, 3987, 3990, 3992, 4001, 4027, 4029, 4072,

165 4194, 4372, 4539, 4543, 4584, 4610, 4620, 4626, 4632, 4641, 4657,

166 4669, 4673, 4674, 4686, 4688, 4690, 4691, 4698, 4709, 4715, 4734,

167 4736, 4746, 4755, 4777, 4778, 4783, 4784, 4789, 4792, 4804, 4812,

168 4814, 4824, 4825, 4830, 4835, 4840, 4841, 4851, 4856, 4858, 4862,

169 4869, 4890, 4891, 4897, 4901, 4903, 4906, 4914

170 <223> OTHER INFORMATION: n = a, t, g or c

172 <400> SEQUENCE: 5

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177	agtcaggccg	ggacgttagc	gcctgcgcgg	gacctcact	tgccaccaag	rmccccacaa	300
178	accccgcccc	atcctgyctt	acgccccgcc	ccaaggtcgg	ttctccccga	cccgggggtc	360
W--> 179	ccgcccccaa	ggnccgtcct	ccccgcccc	gccggttggt	ggcgggcgca	tgctgccccg	420
180	atataaagg	tcggccccac	atcccaggga	ccagcgagcg	gccttgagag	gctctggctc	480
181	ttgcttctta	ggcgggccga	ggacgcatg	gccgagtgc	cgacactcgg	ggaggcagtc	540
182	accgaccacc	cggaccgcct	gtgggcctgg	gagaagttcg	tgtatttga	cgagaagcag	600
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186	gatgaatagg	agttcctcca	ggcagggaag	aagggtgga	aaacccccaa	ggaatgtcgg	840
187	tcaaaggggt	ggaccagtg	cctgtggagt	gtgactataa	tggtgactac	agcaggcatt	900
188	ttctgggctt	cggggtccta	atccttaaaa	atgggtatct	ctaagtgact	catccatatg	960
189	gccgattatc	ggaatcatct	caggtgggtc	ccagaaatct	gtatttttaa	aaagaacccw	1020
190	cmacagttta	gggtccaacc	caggcataac	caaaacactg	gcctaagagt	tgtgaagtat	1080
191	tttccacact	accctctggg	ctttatttaa	gamaaccaa	tttaacaagt	gatgtcgtag	1140
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196	agtgttgayt	gtgtctctac	agatgggaaa	ctgaggcaca	cmaaatgtac	atttgtccga	1440
197	ggtaagattg	ctagtaggta	atgggggttg	aattctaggc	tcttaaccac	cacaaaatct	1500
198	gcatttttat	tggcatttca	atttttttaa	tatgttttta	ctttaaaaaa	caagttaaat	1560
199	acttactttt	ttaaaatcaa	aatttgaa	aataatttga	agattcagtg	gatttctttt	1620
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201	tttttcataa	aatcatgaaa	catgcycmc	maaaaaataa	ccactasca	actgtgggac	1740
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203	gttttttttt	tttctgtag	awacagggtc	tcgctctgtg	acccaggctg	gtctyaaact	1860
204	cctggcctca	agcgatcctc	ctgcctctgc	cttccaaagc	actggaatta	caagtgtgag	1920
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218	tacctgtgtg	ttccagctag	gggaagscsc	aggagtggag	aatggaggga	gtggagggct	2760
219	ctggccgata	aatgccttct	ctctctctct	gcctctcaga	ttgacggcgt	ggaggacatg	2820
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221	ttagtccaca	gtggctgtat	cagaaagaaa	gaccacccct	tctccatgaa	ggcagtgtcta	2940
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226 gcaatctcag ctcactgaaa gctctgcctc ccgggttcac accattctcc tgcctcagcc 3240
227 ctcgagtag ctgggactac aggcgcccgc caccacacct ggctaatttt tttttttttw 3300
228 ttwtwttttt tagtagagmc ggggtttcac cgtgttagcc aggatggtct cgatctcctg 3360
229 acctcatgat ctgcccgcct cggcctccca aagtgcctggg attacaggca tgagccacca 3420
230 cgtccggcct taccattgct ttattaaata agcactgggtg cttgattata tcagctgagc 3480
231 cagatattag atacgctatt gaggtttgrg gaaataagag taccaaaact cagaaatgag 3540
232 ttgaagtata gtgacatctt cagattacag acccagggtg cagaatttgc cttggctcag 3600
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234 ccacccctcg scactgcctg gtcctttcct tcacccttga ttctgtcttc tttgtcctt 3720
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237 ctgggggtctg ggaggaatgg acagacagag gatgagctct acccrgggcc tgsaggacct 3900
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243 ctgctgckk gtraacacgm sksyrmacag stgmswrccc gtaaacacgc ctgcaaacgc 4260
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255 ct 4922
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259 <212> TYPE: DNA
C--> 260 <213> ORGANISM: Artificial
262 <220> FEATURE:
263 <223> OTHER INFORMATION: Description of Artificial Sequence: p9A primer
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268 <210> SEQ ID NO: 7
269 <211> LENGTH: 20
270 <212> TYPE: DNA
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273 <220> FEATURE:
274 <223> OTHER INFORMATION: Description of Artificial Sequence: Rev III primer
276 <400> SEQUENCE: 7
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279 <210> SEQ ID NO: 8

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Use of n and / or Xaa has been detected in the Sequence Listing. Review the Sequence Listing to ensure a corresponding explanation is present in the <220> to <223> fields of each sequence using n or Xaa.

VERIFICATION SUMMARY

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L:109 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:3
L:129 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:4
L:151 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:156 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:5
L:161 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:5
L:179 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5
L:184 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5
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L:326 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:12
L:350 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12